

1653

P#9

RAW SEQUENCE LISTING DATE: 02/22/2001
 PATENT APPLICATION: US/09/446,543A TIME: 09:07:20

Input Set : A:\2472USOP.txt
 Output Set: N:\CRF3\02222001\I446543A.raw

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see p. 5

3 <110> APPLICANT: HINUMA, Shuji
 4 KAWAMATA, Yuji
 5 FUJII, Ryo
 6 MATSUMOTO, Hirokazu
 8 <120> TITLE OF INVENTION: Prolactin Secretion Modulator
 10 <130> FILE REFERENCE: 2472USOP
 12 <140> CURRENT APPLICATION NUMBER: US 09/446,543A
 13 <141> CURRENT FILING DATE: 1999-12-20
 15 <150> PRIOR APPLICATION NUMBER: PCT/JP98/02765
 16 <151> PRIOR FILING DATE: 1998-06-22
 18 <150> PRIOR APPLICATION NUMBER: JP 9-165437
 19 <151> PRIOR FILING DATE: 1997-06-23
 21 <160> NUMBER OF SEQ ID NOS: 99
 23 <170> SOFTWARE: PatentIn version 3.0
 25 <210> SEQ ID NO: 1
 26 <211> LENGTH: 98
 27 <212> TYPE: PRT
 28 <213> ORGANISM: Bovine
 30 <400> SEQUENCE: 1
 32 Met Lys Ala Val Gly Ala Trp Leu Leu Cys Leu Leu Leu Gly Leu
 33 1 5 10 15
 35 Ala Leu Gln Gly Ala Ala Ser Arg Ala His Gln His Ser Met Glu Ile
 36 20 25 30
 38 Arg Thr Pro Asp Ile Asn Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg
 39 35 40 45
 41 Pro Val Gly Arg Phe Gly Arg Arg Arg Ala Ala Pro Gly Asp Gly Pro
 42 50 55 60
 44 Arg Pro Gly Pro Arg Arg Val Pro Ala Cys Phe Arg Leu Glu Gly Gly
 45 65 70 75 80
 47 Ala Glu Pro Ser Arg Ala Leu Pro Gly Arg Leu Thr Ala Gln Leu Val
 48 85 90 95
 50 Gln Glu
 53 <210> SEQ ID NO: 2
 54 <211> LENGTH: 294
 55 <212> TYPE: DNA
 56 <213> ORGANISM: bovine
 58 <400> SEQUENCE: 2
 59 atgaaggcgg tgggggcctg gctcctctgc ctgctgctgc tgggcctggc cctgcagggg 60
 61 gctgccagca gaggccacca gcactccatg gagatccgca ccccgacat caacctgac 120
 63 tggtaacgcr ggcgtgggat ccggcccgta ggccgcttcg gccggcgaa agctgcccyg 180
 65 ggggacggac ccaggcctgg ccccccggct gtgccggcct gcttccgct ggaaggcggg 240
 67 gctgagcccl cccgagcccl cccggggcgg ctgacggccc agctggloca ggaa 294
 70 <210> SEQ ID NO: 3
 71 <211> LENGTH: 29
 72 <212> TYPE: PRT
 73 <213> ORGANISM: artificial
 75 <220> FEATURE:

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76 <223> OTHER INFORMATION: bovine fragment (23-51)
78 <400> SEQUENCE: 3
80 Ser Arg Ala His Gln His Ser Met Glu Ile Arg Thr Pro Asp Ile Asn
81 1 5 10 15
83 Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro Val Gly
84 20 25
86 <210> SEQ ID NO: 4
87 <211> LENGTH: 19
88 <212> TYPE: PRT
89 <213> ORGANISM: artificial
91 <220> FEATURE:
92 <223> OTHER INFORMATION: bovine fragment (34-52)
94 <400> SEQUENCE: 4
96 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro
97 1 5 10 15
99 Val Gly Arg
102 <210> SEQ ID NO: 5
103 <211> LENGTH: 31
104 <212> TYPE: PRT
105 <213> ORGANISM: artificial
107 <220> FEATURE:
108 <223> OTHER INFORMATION: bovine fragment (23-53)
110 <400> SEQUENCE: 5
112 Ser Arg Ala His Gln His Ser Met Glu Ile Arg Thr Pro Asp Ile Asn
113 1 5 10 15
115 Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro Val Gly Arg Phe
116 20 25 30
118 <210> SEQ ID NO: 6
119 <211> LENGTH: 32
120 <212> TYPE: PRT
121 <213> ORGANISM: artificial
123 <220> FEATURE:
124 <223> OTHER INFORMATION: bovine fragment (23-54)
126 <400> SEQUENCE: 6
128 Ser Arg Ala His Gln His Ser Met Glu Ile Arg Thr Pro Asp Ile Asn
129 1 5 10 15
131 Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro Val Gly Arg Phe Gly
132 20 25 30
134 <210> SEQ ID NO: 7
135 <211> LENGTH: 33
136 <212> TYPE: PRT
137 <213> ORGANISM: artificial
139 <220> FEATURE:
140 <223> OTHER INFORMATION: bovine fragment (23-55)
142 <400> SEQUENCE: 7
144 Ser Arg Ala His Gln His Ser Met Glu Ile Arg Thr Pro Asp Ile Asn
145 1 5 10 15
147 Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro Val Gly Arg Phe Gly
148 20 25 30

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150 Arg
153 <210> SEQ ID NO: 8
154 <211> LENGTH: 20
155 <212> TYPE: PRT
156 <213> ORGANISM: artificial
158 <220> FEATURE:
159 <223> OTHER INFORMATION: bovine fragment (34-53)
161 <400> SEQUENCE: 8
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164 1 5 10 15
166 Val Gly Arg Phe
167 20
169 <210> SEQ ID NO: 9
170 <211> LENGTH: 21
171 <212> TYPE: PRT
172 <213> ORGANISM: artificial
174 <220> FEATURE:
175 <223> OTHER INFORMATION: bovine fragment (34-54)
177 <400> SEQUENCE: 9
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180 1 5 10 15
182 Val Gly Arg Phe Gly
183 20
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186 <211> LENGTH: 22
187 <212> TYPE: PRT
188 <213> ORGANISM: artificial
190 <220> FEATURE:
191 <223> OTHER INFORMATION: bovine fragment (34-55)
193 <400> SEQUENCE: 10
195 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro
196 1 5 10 15
198 Val Gly Arg Phe Gly Arg
199 20
201 <210> SEQ ID NO: 11
202 <211> LENGTH: 87
203 <212> TYPE: DNA
204 <213> ORGANISM: bovine
206 <400> SEQUENCE: 11
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209 gcrggccgtg ggateccggcc cgtgggc 87
212 <210> SEQ ID NO: 12
213 <211> LENGTH: 57
214 <212> TYPE: DNA
215 <213> ORGANISM: bovine
217 <400> SEQUENCE: 12
218 acccccgaca tcaacctgc ctggtacgcr ggccgtggga tccggcccgt gggccgc 57
221 <210> SEQ ID NO: 13
222 <211> LENGTH: 93

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223 <212> TYPE: DNA
224 <213> ORGANISM: bovine
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229 gcrqgcctgt ggcaccggcc cgtgggcccgc ttc 93
232 <210> SEQ ID NO: 14
233 <211> LENGTH: 96
234 <212> TYPE: DNA
235 <213> ORGANISM: bovine
237 <400> SEQUENCE: 14
238 agcagagccc accagcactc catggagatc cgcacccccg acatcaaccc tgcctgggtac 60
240 gcrqgcctgt ggcaccggcc cgtgggcccgc ttcggc 96
243 <210> SEQ ID NO: 15
244 <211> LENGTH: 99
245 <212> TYPE: DNA
246 <213> ORGANISM: bovine
248 <400> SEQUENCE: 15
249 agcagagccc accagcactc catggagatc cgcacccccg acatcaaccc tgcctgggtac 60
251 gcrqgcctgt ggcaccggcc cgtgggcccgc ttcggcgg 99
254 <210> SEQ ID NO: 16
255 <211> LENGTH: 60
256 <212> TYPE: DNA
257 <213> ORGANISM: bovine
259 <400> SEQUENCE: 16
260 acccccagaca tcaaccctgc ctggtacgcr ggcgctggga tccggcccgt gggccgcttc 60
263 <210> SEQ ID NO: 17
264 <211> LENGTH: 63
265 <212> TYPE: DNA
266 <213> ORGANISM: bovine
268 <400> SEQUENCE: 17
269 acccccagaca tcaaccctgc ctggtacgcr ggcgctggga tccggcccgt gggccgcttc 60
271 ggc 63
274 <210> SEQ ID NO: 18
275 <211> LENGTH: 66
276 <212> TYPE: DNA
277 <213> ORGANISM: bovine
279 <400> SEQUENCE: 18
280 acccccagaca tcaaccctgc ctggtacgcr ggcgctggga tccggcccgt gggccgcttc 60
282 ggcgg 66
285 <210> SEQ ID NO: 19
286 <211> LENGTH: 91
287 <212> TYPE: PRT
288 <213> ORGANISM: Human
290 <400> SEQUENCE: 19
292 Leu Val Leu Val Ile Ala Arg Val Arg Arg Leu His Asn Val Thr Asn
293 1 5 10 15
295 Phe Leu Ile Gly Asn Leu Ala Leu Ser Asp Val Leu Met Cys Thr Ala
296 20 25 30
298 Cys Val Pro Leu Thr Leu Ala Tyr Ala Phe Glu Pro Arg Gly Trp Val

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Input Set : A:\2472US0P.txt
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299          35          40          45
301 Phe Gly Gly Gly Leu Cys His Leu Val Phe Phe Leu Gln Pro Val Thr
302          50          55          60
304 Val Tyr Val Ser Val Phe Thr Leu Thr Thr Ile Ala Val Asp Arg Tyr
305 65          70          75          80
307 Val Val Leu Val His Pro Leu Arg Arg Arg Ile
308          85          90
310 <210> SEQ ID NO: 20
311 <211> LENGTH: 59
312 <212> TYPE: PRT
313 <213> ORGANISM: human
315 <400> SEQUENCE: 20
317 Gly Leu Leu Leu Val Thr Tyr Leu Leu Pro Leu Leu Val Ile Leu Leu
318 1          5          10          15
320 Ser Tyr Val Arg Val Ser Val Lys Leu Arg Asn Arg Val Val Pro Gly
321          20          25          30
323 Cys Val Thr Gln Ser Gln Ala Asp Trp Asp Arg Ala Arg Arg Arg Arg
324          35          40          45
326 Thr Phe Cys Leu Leu Val Val Val Val Val
327          50          55
329 <210> SEQ ID NO: 21
330 <211> LENGTH: 370
331 <212> TYPE: PRT
332 <213> ORGANISM: human
334 <400> SEQUENCE: 21
336 Met Ala Ser Ser Thr Thr Arg Gly Pro Arg Val Ser Asp Leu Phe Ser
337 1          5          10          15
339 Gly Leu Pro Pro Ala Val Thr Thr Pro Ala Asn Gln Ser Ala Glu Ala
340          20          25          30
342 Ser Ala Gly Asn Gly Ser Val Ala Gly Ala Asp Ala Pro Ala Val Thr
343          35          40          45
345 Pro Phe Gln Ser Leu Gln Leu Val His Gln Leu Lys Gly Leu Ile Val
346          50          55          60
348 Leu Leu Tyr Ser Val Val Val Val Gly Leu Val Gly Asn Cys Leu
349 65          70          75          80
351 Leu Val Leu Val Ile Ala Arg Val Arg Arg Leu His Asn Val Thr Asn
352          85          90          95
354 Phe Leu Ile Gly Asn Leu Ala Leu Ser Asp Val Leu Met Cys Thr Ala
355          100          105          110
357 Cys Val Pro Leu Thr Leu Ala Tyr Ala Phe Glu Pro Arg Gly Trp Val
358          115          120          125
360 Phe Gly Gly Gly Leu Cys His Leu Val Phe Phe Leu Gln Pro Val Thr
361          130          135          140
363 Val Tyr Val Ser Val Phe Thr Leu Thr Thr Ile Ala Val Asp Arg Tyr
364 145          150          155          160
366 Val Val Leu Val His Pro Leu Arg Arg Arg Ile Ser Leu Arg Leu Ser
367          165          170          175
369 Ala Tyr Ala Val Leu Ala Ile Trp Ala Leu Ser Ala Val Leu Ala Leu
370          180          185          190

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F.4.1.

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY DATE: 02/22/2001
PATENT APPLICATION: US/09/446,543A TIME: 09:07:21

Input Set : A:\2472US0P.txt
Output Set: N:\CRF3\02222001\I446543A.raw

L:625 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:649 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:661 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:31
L:667 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:31
L:673 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:685 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:32
L:691 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:733 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:769 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36
L:793 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37
L:1337 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73
L:1340 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73
L:1371 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74